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FITTLE OF INVENTION: APOPTOSIS MODULATOR BCL-B
FILT REFERENCE: 8014-014-US
CURRENT APPLICATION NUMBER: US/10/071,174A
CURRENT FILING DATE: 2002-02-07
PRIOR APPLICATION NUMBER: 60/267,166
PRIOR PILING DATE: 2001-02-07
NUMBER OF SEQ ID NOS: 37
SOPTWARE: PatentIn version 3.3
SEQ ID NO 1
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US-10-071-174A-1-COPY
; Sequence 1, Application US/10071174A
; GENERAL INFORMATION:
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APPLICANT: Reed, John C.
APPLICANT: Ke, Ning
APPLICANT: Adam, Godzik
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LENGTH: 887
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ORGANISM: Homo sapiens
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| Sequence 37-Application US/10071174A
| GENERAL INFORMATION:
| APPLICANT: The Burnham Institute
| APPLICANT: The Arch Institute
APPLICANT: The Burnham Institute
APPLICANT: Reed, John C.
APPLICANT: Ke, Ning
APPLICANT: Ke, Ning
APPLICANT: Adam, Godzik
TITLE OF INVENTION: APOPTOSIS MODULATOR BCL-B AND METHODS FOR MAKING AND USING
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US-10-071-174A-1-COPY

Query Match

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US-10-071-174A-37-COPY
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Best Local Similarity
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FILE REFERENCE: 8014-014-US
CURRENT APPLICATION NUMBER: US/10/071,174A
CURRENT FILING DATE: 2002-02-07
PRIOR APPLICATION NUMBER: 60/267,166
PRIOR FILING DATE: 2001-02-07
NUMBER OF SEQ ID NOS: 37
                                                                                                                                                                                                                                                                                                                                                                    CURRENT APPLICATION NUMBER: US/10/071,174A
CURRENT FILING DATE: 2002-02-07
PRIOR APPLICATION NUMBER: 60/267,166
PRIOR FILING DATE: 2001-02-07
                                                                                                                                                                                                                                                                                                                                                                                                                                  TITLE OF INVENTION: APOPTOSIS MODULATOR BCL-B AND METHODS FOR MAKING AND USING THE TITLE OF INVENTION: SAME FILE REFERENCE: 8014-014-US
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ORGANISM: Homo sapiens
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Ke, Ning
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; TYPE: DNA
; ORGANISM: Homo sapiens
; FEATURE:
; NAME/KEY: misc_feature
; LOCATION: (94)...(94)
; OTHER INFORMATION: n is a
US-10-071-174A-37-COPY
Title:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: The Burnham Institute APPLICANT: Reed, John C.
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SEQ ID NO 37
LENGTH: 130
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                                                                                                                 Matches
                                                                                                                                               Query Match
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APPLICANT: Adam, Godzik
TITLE OF INVENTION: APOPTOSIS MODULATOR BCL-B AND METHODS FOR MAKING AND USING THE
TITLE OF INVENTION: SAME
FILE REFERENCE: 8014-014-US
CURRENT PEPLICATION NUMBER: US/10/071,174A
CURRENT FILING DATE: 2002-02-07
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PRIOR APPLICATION NUMBER: 60/267,166
PRIOR FILING DATE: 2001-02-07
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APPLICANT: Reed, John C.
APPLICANT: Ke, Ning
APPLICANT: Adam, Godzik
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                                                                                                                                                                                                                                                          TYPE: DNA
ORGANISM: Homo sapiens
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APPLICANT: Adam, Godzik
APPLICANT: Adam, Godzik
TITLE OF INVENTION: APOPTOSIS MODULATOR BCL-B AN
TITLE OF INVENTION: SAME
FILE REFERENCE: 8014-0.14-US
CURRENT APPLICATION UMBER: US/10/071,174A
CURRENT FILING DATE: 2002-02-07
PRIOR APPLICATION NUMBER: 60/267,166
PRIOR FILING DATE: 2001-02-07
NUMBER OF SEQ ID NOS: 37
                                                                                                                    SOFTWARE: PatentIn version 3.3 SEQ ID NO 37 LENGTH: 130
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APPLICANT: The Burnham
APPLICANT:
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TYPE: DNA
ORGANISM: Homo sapiens
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                              NAME/KEY: misc_feature
LOCATION: (94)..(94)
                                                                                                         TYPE: DNA
                                                                                    ORGANISM: Homo sapiens
                                                                 FEATURE:
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RESULT 4
US-10-071-174A-1-COPY/c
Sequence 1, Application US/10071174A
GENERAL INFORMATION:
APPLICANT: The Burnham Institute
APPLICANT: Reed, John C.
APPLICANT: Ke, Ning
APPLICANT: Adam, Godzik
TITLE OF INVENTION: APOPTOSIS MODULATOR BCL-B AND METHODS FOR MAKING AND USING THE
TITLE OF INVENTION: SAME
FILE REFERENCE: 8014-014-US
CURRENT APPLICATION NUMBER: US/10/071,174A
CURRENT FILING DATE: 2002-02-07
PRIOR APPLICATION NUMBER: 60/267,166
PRIOR FILING DATE: 2001-02-07
NUMBER OF SEQ ID NOS: 37
SOFTWARE: PATENTIN VERSION 3.3
SEQ ID NO 1
LENGTH: 887
TYPE: DNA
ORGANISM: Homo sapiens
US-10-071-174A-1-COPY
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Search completed: June 6, 2005, 09:08:24 Job time: 0.127827 secs
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                                                                                                                                                                                                                                                                                     Query Match 17.2%; Score 22.4; DB 1; Length 887; Best Local Similarity 58.5%; Pred. No. 0; Matches 38; Conservative 0; Mismatches 27; Indels 0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match 19.4%; Score 25.2; DB 1; Length 130; Best Local Similarity 58.3%; Pred. No. 0; Matches 42; Conservative 0; Mismatches 30; Indels 0;
                                                                                             271 GCGGT 267
                                                                                                                                                                                               331 CCAGGTGGGGCCGGGGCTGTCGGAGAGCACGGAATCCGCCATCAGCGCCACCAGCTCGAA 272
                                                                                                                                                                                                                      33 CGAGCTGGTGGCGCTGATGGCGGATTCCGTGCTCTCCGACAGCCCCGGCCCCACCTGGGA 92
                                                                                                                                             93 GNAGT 97
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                86 CCTGGGAGNAGT 97
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 26 ACCGCTTCGAGCTGGTGGCGCTGATGGCGGATTCCGTGCTCTCCGACAGCCCCGGCCCCA 85
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                37 GCTCGAÁGCGGT 26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                97 ACTNOTOCCAGGTGGGGCCGGGGCTGTCGGAGAGCACGGAATCCGCCATCAGCGCCACCA 38
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                          0;
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